Please amend the specification as follows.

On page 18, please replace the first full paragraph, which begins on line 4 with

"Specific forward" with the following amended paragraph:

Specific forward (STGROF1) and reverse primers (STGROR2) were designed to

be capable of amplifying groEL2 gene fragment in all Streptomyces species. S. lividans

(GenBank No. X95971) and S. albus (GenBank No. M76658), whose full-length

nucleotide sequences of groEL2 gene were already sequenced for other purposes, and

T. paurometabola (GenBank No. AF35257 AF352578), which belongs to Tsukamurella

species closely related to Streptomyces species in a phylogenetic tree, were subjected

to sequencing analysis, and forward primer STGROF1 (5'-CCATCGCCAAGGAGAT

CGAGCT-3': SEQ ID NO: 1) and reverse primer STGROR2 (5'-

TGAAGGTGCCRCGGATCTTGTT-3': SEQ ID NO: 2) that are capable of specifically

amplifying all Streptomyces species were prepared therefrom (Fig. 1). The primer pair

of STGROF1 and STGROR2 is new which has not been used previously for amplifying

Streptomyces species.

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